

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/832,189

DATE: 05/07/2001
TIME: 12:09:30

Input Set : A:\Pto.amc
Output Set: N:\CRF3\05072001\I832189.raw

2 <110> APPLICANT: RIKEN
4 <120> TITLE OF INVENTION: Truncated Reelin Protein and DNA Encoding the Same
6 <130> FILE REFERENCE: PH-1167
C--> 8 <140> CURRENT APPLICATION NUMBER: US/09/832,189
C--> 9 <141> CURRENT FILING DATE: 2001-04-11
11 <150> PRIOR APPLICATION NUMBER: JP 2000-109954
12 <151> PRIOR FILING DATE: 2000-04-11
14 <160> NUMBER OF SEQ ID NOS: 28
16 <170> SOFTWARE: PatentIn Ver. 2.0
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 2274
20 <212> TYPE: DNA
21 <213> ORGANISM: Xenopus laevis
23 <220> FEATURE:
24 <221> NAME/KEY: CDS
25 <222> LOCATION: (157)..(1455)
27 <220> FEATURE:
28 <221> NAME/KEY: sig_peptide
29 <222> LOCATION: (157)..(234)
31 <220> FEATURE:
32 <221> NAME/KEY: misc_feature
33 <222> LOCATION: (241)..(726)
34 <223> OTHER INFORMATION: F-spondin domain
36 <220> FEATURE:
37 <221> NAME/KEY: misc_feature
38 <222> LOCATION: (847)..(1197)
39 <223> OTHER INFORMATION: CR-50 epitope region
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46 atgaagaaag ctcattaaga acagtggac ccagga atg gaa ctg ctc cac acc 174
47 Met Glu Leu Leu His Thr
48 1 5
50 ttc tgc ggt ggg cgc tgg act ttg ctg ctc acg ggg atc ttg tgc 222
51 Phe Cys Gly Gly Arg Trp Thr Leu Leu Leu Phe Thr Gly Ile Leu Cys
52 10 15 20
54 ttt gtt gtt gcc cgc gga gtg ggg tat tat ccc agg ttc tct cca ttc 270
55 Phe Val Val Ala Arg Gly Val Gly Tyr Tyr Pro Arg Phe Ser Pro Phe
56 25 30 35
58 ttt ttc ctt tgc act cat cat gga gaa ctg gaa gga gat ggg gaa caa 318
59 Phe Phe Leu Cys Thr His His Gly Glu Leu Glu Gly Asp Gly Glu Gln
60 40 45 50
62 gga gaa gtg ctc atc tct ctg cac ctg gcg ggc aac ccc agc tac tac 366
63 Gly Glu Val Leu Ile Ser Leu His Ala Gly Asn Pro Ser Tyr Tyr
64 55 60 65 70
66 ata cct ggg cag gag tac cat gtg acc ata tcc act agt acc ttc ttt 414
67 Ile Pro Gly Gln Glu Tyr His Val Thr Ile Ser Thr Ser Thr Phe Phe

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see item 10
on Error
Summary Sheet

FYI

Please Note:

Use f n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.